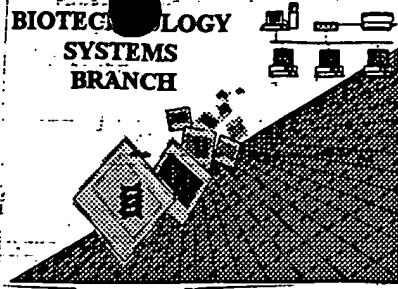


0500

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/721,543

Source: O/PE

Date Processed by STIC: 12/7/2000

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/721,543

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 Wrapped Nucleic The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".

2 Wrapped Aminos The amino acid-number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".

3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.

4 Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.

5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.

6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.

7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

8 Skipped Sequences (OLD RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s)

9 Skipped Sequences (NEW RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence
<210> sequence id number
<400> sequence id number
000

10 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents

11 Use of <213>Organism (NEW RULES) Sequence(s) are missing this mandatory field or its response.

12 Use of <220>Feature (NEW RULES) Sequence(s) are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec 1 823 of new Rules)

13 PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other means to copy file to floppy disk.

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and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

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- GRAY SCALE DOCUMENTS

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As rescanning documents *will not* correct images,
please do not report the images to the
Image Problem Mailbox.

Q1 PE

RAW SECTION LISTING
US/09/121,543
A:SEARCH.LST
H:\NCB\12072000\121543.raw

Does Not Comply
Corrected Diskette Needed

OK 13 <140> CURRENT APPLICATION NUMBER: US793721,543

pp-1.5

per 1.823 of new Sequence Rules, the only valid
responses are: Unknown, Artificial Sequence, or
Scientific name
(genus/species)
(see circled
portion of item 12
on Error Summary
sheet)

RAY SHOTLOCK LISTING
10-10-2000, 04:00
At: REQLIST.txt
At: C:\CRF3\12072000\1721543.raw

RAM SEQUENCE LISTING

Ergonomics in Design 2011, 20(1)

At: **SEGOLIST.LST**

File: C:\12072000\1721540.raw

1. 100 2. 100 3. 100 4. 100 5. 100 6. 100 7. 100 8. 100 9. 100 10. 100 11. 100 12. 100 13. 100 14. 100 15. 100 16. 100 17. 100 18. 100 19. 100 20. 100 21. 100 22. 100 23. 100 24. 100 25. 100 26. 100 27. 100 28. 100 29. 100 30. 100 31. 100 32. 100 33. 100 34. 100 35. 100 36. 100 37. 100 38. 100 39. 100 40. 100 41. 100 42. 100 43. 100 44. 100 45. 100 46. 100 47. 100 48. 100 49. 100 50. 100 51. 100 52. 100 53. 100 54. 100 55. 100 56. 100 57. 100 58. 100 59. 100 60. 100 61. 100 62. 100 63. 100 64. 100 65. 100 66. 100 67. 100 68. 100 69. 100 70. 100 71. 100 72. 100 73. 100 74. 100 75. 100 76. 100 77. 100 78. 100 79. 100 80. 100 81. 100 82. 100 83. 100 84. 100 85. 100 86. 100 87. 100 88. 100 89. 100 90. 100 91. 100 92. 100 93. 100 94. 100 95. 100 96. 100 97. 100 98. 100 99. 100 100. 100

LAW OFFICES OF R. L. ST. LUCE

A: SEDLIST.txt
B: C:\12072009\1721541.zaw

5

OK

FYI:

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

1947. The current and former Home Office Ministers Replaced "Accrued" by "Accrued and
Paid" in 1947, and "Accrued" by "Paid" again for SMC 1949.
The "Accrued" period is "Paid" again for SMC 1950.
The "Accrued" period is "Paid" again for SMC 1951.
The "Accrued" period is "Paid" again for SMC 1952.
The "Accrued" period is "Paid" again for SMC 1953.
The "Accrued" period is "Paid" again for SMC 1954.